

# *Toward multifactorial null models of range contraction in terrestrial vertebrates*

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*List of authors:* Pablo M. Lucas<sup>1</sup>, Manuela González-Suárez<sup>1,2</sup>, Eloy Revilla<sup>1</sup>

*Institute of origin:*

<sup>1</sup>Department of Conservation Biology, Estación Biológica de Doñana EBD-CSIC

Calle Americo Vespucio s/n 41092 Seville Spain

<sup>2</sup>Ecology and Evolutionary Biology, School of Biological Sciences, University of

Reading, Whiteknights, Reading RG6 6AS, UK

*Corresponding author:* PML, Tel: (+34) 954 466700 (Ext. 1125), Fax: (+34) 954

621125, lucas.pablo.2010@gmail.com

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## ABSTRACT

The contraction of a species' distribution range, which results from the extirpation of local populations, generally precedes its extinction. Therefore, understanding drivers of range contraction is important for conservation and management. Although there are many processes that can potentially lead to local extirpation and range contraction, three main null models have been proposed: demographic, contagion, and refuge. The first two models postulate that the probability of local extirpation for a given area depends on its relative position within the range; but these models generate distinct spatial predictions because they assume either a ubiquitous (demographic) or a clinal (contagion) distribution of threats. The third model (refuge) postulates that extirpations are determined by the intensity of human impacts, leading to heterogeneous spatial predictions potentially compatible with those made by the other two null models. A few previous studies have explored the generality of some of these null models, but we present here the first comprehensive evaluation of all three models. Using descriptive indices and regression analyses we contrast the predictions made by each of the null models using empirical spatial data describing range contraction in 386 terrestrial vertebrates (mammals, birds, amphibians, and reptiles) distributed across the World. Observed contraction patterns do not consistently conform to the predictions of any of the three models, suggesting that these may not be adequate null models to evaluate range contraction dynamics among terrestrial vertebrates. Instead, our results support alternative null models that account for both relative position and intensity of human impacts. These new models provide a better multifactorial baseline to describe range contraction patterns in vertebrates. This general baseline can be used to explore how additional factors influence contraction, and ultimately extinction for particular areas or species as well as to predict future changes in light of current and new threats.

38    *Keywords:* Border, extinction, habitat loss, historical range, human, land use, range  
39    dynamics.

## INTRODUCTION

Species extinctions generally start with the vanishing of particular populations that continue until no populations remain (Yackulic et al. 2011). In other words, complete extinction is usually preceded by a contraction of the distribution range that results from the extirpation of local populations. Local extirpations and contractions are considered good descriptors of biological capital loss, possibly even preferable to quantifying extinction itself (Ceballos and Ehrlich 2002, Davis et al. 1998). Therefore, understanding the general dynamics of range contraction is key for effective conservation (Safi and Pettorelli 2010). The list of proximate and ultimate causes of local extinction is long, and taxon-dependent (Cahill et al. 2012, González-Suárez and Revilla 2014); thus, we may expect a wide variety of range contraction patterns. Nevertheless, ecologists and conservation biologists have used null models or simple hypotheses to describe the expected spatial patterns of local extinction and range contraction, especially when detailed information is not available.

Null models are representations based on the simplest and most general mechanisms, and deliberately focus on a few key factors or processes to provide a baseline for comparison with empirical observations or with more complex models (Gotelli 2001). The simplicity of null models can be useful for species for which little information exists, as well as in theoretical studies (Hanski 1998, Hanski and Ovaskainen 2000). Generalized patterns of distribution range contraction have been described in the literature using three different null models: demographic, contagion, and refuge. These models describe contraction based on distinct mechanisms derived from theoretical principles in ecology, biogeography, and conservation biology (Hanski 1998, Hemerik et al. 2006); and have been used in empirical studies as baselines to determine the role of additional factors or to broadly describe observed contraction

patterns (Franco et al. 2006, Parmesan 1996, Pomara et al. 2014, Thomas et al. 2004, Turvey et al. 2015, Yackulic et al. 2011).

The demographic null model derives from basic population dynamic principles, and from the ecological assumption which postulate that environmental conditions and resources at the center of a distribution range are more suitable than at the border, resulting in higher population growth rates and thus, higher abundance in central areas (Brown 1995, Lawton 1993). Because extinction is directly determined by population abundance (Brown 1971, David et al. 2003, Jones and Diamond 1976, Pimm et al. 1988), when the drivers of extinction (threats) are ubiquitous, central areas would have lower extinction/extirpation risk (MacArthur and Wilson 1967). Assuming threats are indeed ubiquitous, this null model then predicts that populations would be first extirpated along the historical border (where density is lower) and would continue toward the center, where the last (most dense) population would be found (Fig. 1). The contagion null model, on the other hand, assumes that the threats have clinal distribution, with threats spreading across the landscape with distinct directionality, like a contagious disease (Channell and Lomolino 2000a, Channell and Lomolino 2000b, Lawton 1993). Based on this clinal threat pattern, the contagion null model predicts that populations would be first extirpated in the historical border closest to the extinction driver's origin, and then as the threat spreads across the range, the central areas would become extirpated until only the historical border located farthest from the initial point remains (Fig. 1). Finally, the refuge model assumes that more humanized land uses are associated with higher risk of extinction (Ceballos and Ehrlich 2002, Fisher 2011, Hoffmann et al. 2010, Laliberte and Ripple 2004, Li et al. 2015, Pomara et al. 2014, Schipper et al. 2008, Yackulic et al. 2011), and predicts that populations would be first extirpated in areas that are more modified and heavily used by humans. According to

this model, the last population will be located in the least used area, which represents a final refuge for the species (Fig. 1).

Some of the assumptions and the predictions of primarily the demographic and contagion models have been tested by previous studies, which collectively suggest these models may not be broadly applicable (Ceballos and Ehrlich 2002, Fisher 2011, Hemerik et al. 2006, Laliberte and Ripple 2004, Sagarin and Gaines 2002, Thomas et al. 2008, Yackulic et al. 2011). However, there has been no comprehensive evaluation of all three null models; partly because spatial data quantifying range contraction at the global scale are limited, but also because there are important methodological challenges including the difficulties in defining a unique center and a relative position within a species range. In this study we overcome these challenges to simultaneously evaluate these three null models using a global dataset for 386 terrestrial vertebrates (mammals, birds, amphibians and reptiles). We first identify the key predictions derived from each null model and then, using descriptive indices and regression analyses, we evaluate if empirical range contraction data conform to the models predictions. Our goals are: 1) to determine which, if any, of the proposed null models represents the most adequate general baseline to explain range contractions; 2) if necessary, to propose and evaluate alternative multifactorial null models; and 3) to provide a more consistent framework regarding the general underlying causes of range contraction dynamics among terrestrial vertebrates.

## **METHODS**

### **Spatial distribution data**

We used global distribution data of 386 terrestrial vertebrates (International Union for Conservation of Nature 2010) with known range contraction (i.e., a distribution with

extirpated areas, where the species was present in the past but is no longer found, and current areas, where the species is currently present, and following the notation of the International Union for Conservation of Nature 2010; detailed information is provided in Appendix 1). Since most species distributions are fragmented and have complex shapes, our analyses were conducted at two different scales. At the range scale, we used data from the complete historical distribution range of each species ( $N=374$ ), which often included multiple fragments separated by unoccupied areas. At the fragment scale, we used data from all individual fragments with observed contraction ( $N=273$ . See Supplementary materials for additional information in data preparation). Supplementary material Appendix 2, Table A3 and A4, and Fig. A1 provide descriptive summaries of these data including total area in  $\text{km}^2$  and percentage of contraction (calculated as the percentage of the historical range area classified as extirpated) for complete ranges and individuals fragments. For complete ranges we also summarize the number of fragments present in the historical, extirpated, and current ranges, as well as the percentage of extirpated fragments (percentage of historical fragments classified as extirpated). Spatial data were projected into an equal area projection (Cylindrical Equal Area) and rasterized.

### **Analyses**

We followed a two-step approach to evaluate the key predictions of each null model (Fig. 1). First, we defined three indices to visually explore the support of model predictions by the empirical data. Second, we defined and compared three regression models that estimate the probability of extirpation based on the key model predictions, thus providing a quantitative test of support for each null model.



## Indexes

The demographic and contagion null models both associate the probability of extirpation with an area's relative position within a range (Fig. 1). Therefore, we defined a position index based on relative distance to the border. We use the border instead of the center because identifying meaningful centers is complicated in complexly shaped and fragmented distributions (Sagarin et al. 2006). For each distribution range and fragment analyzed, we first estimated the geodetic distance from each grid cell to the closest historical border cell (Fig. 2, and see Supplementary material Appendix 2). A geodetic distance is the distance between two unprojected points on the spheroid of the Earth (using the spheroid World Geodetic System 1984, WGS84). Distances were standardized dividing species' values by the maximum distance observed for the range (at range scale) or fragment (at fragment scale) to facilitate comparison among species with different distribution ranges. Using these distance values from each cell to the nearest border, we then calculated the variable *Border* as the arithmetic mean distance to the border from all cells within one area, with *Border\_ext* representing extirpated areas and *Border\_curr* current areas. Using these values we defined the *Centrality Index* =  $Border\_ext / Border\_curr$  for each range and fragment. The demographic null model predicts *Centrality Index* < 1 (extirpated areas are closer to the border), whereas the contagion model predicts *Centrality Index* < 1 only for initial stages of contraction (approximately <50% of the historical range extirpated), and *Centrality Index* > 1 for contractions >50%. Therefore, both the contagion and demographic null models predict the same values of *Centrality Index* in early stages of contraction but different values in later stages. The refuge null model makes no general prediction for the *Centrality Index* (Fig. 1).

The second prediction made by the demographic and contagion null models relates to the directionality in contraction. The demographic null model predicts that contraction occurs in multiple directions, while the contagion null model states that contraction occurs along a unique general direction that can be detected as a predominant contraction angle (Fig. 1). We calculated the geodetic angle of contraction for each extirpated cell as the azimuth of the direction defined by the vector joining each extirpated grid cell with its closest current cell (Fig. 2 and Supplementary material, Appendix 2). Using all angles of contraction for each distribution (complete range or individual fragment) we calculated the *Directionality Index* as the angular concentration. *Directionality Index* ranges from 0 to 1 and is the inverse of the dispersion of the angles (Zar 1999). The demographic null model predicts *Directionality Index* values close to 0 (high angle dispersion) and the contagion null model predicts values close to 1 (a low angle dispersion). The refuge model makes no prediction for the *Directionality Index* (Fig. 1).

The last index we defined captures the predictions of the refuge model (Fig. 1). Although human land use has changed over time and past uses likely influenced observed contraction, data are not available at a global scale to describe past land use. Therefore, we defined land use based on the 1-km resolution MODIS (MCD12Q1) Land Cover Product (Oak Ridge National Laboratory Distributed Active Archive Center 2010). We determined the extent of land classified as covered/used (henceforward used) by humans for each range or fragment (Supplementary material, Appendix 2 and Table A5). From these cell values we then calculated the variables *Land use\_ext* as the proportion of cells used by humans in the extirpated area, and *Land use\_curr* as the proportion of cells used by humans in the current area. Using these variables, we defined a *Land use Index* which is calculated as  $Land\ use\_ext / Land$

*use\_curr*. If extirpated areas have a greater proportion of human use, then *Land use Index* > 1 as predicted by the refuge null model. The contagion and demographic null models make no specific predictions regarding the *Land use Index*. We calculated and investigated the distribution of these three indices for terrestrial vertebrates.

Prior to visualizing the empirical data the behavior of the *Centrality* and *Directionality* indexes was evaluated using simulated scenarios. We sketched three example distribution range areas (Supplementary material Figure A3) for which we simulated two patterns: range contraction towards the center (demographic model), and clinal range contraction (contagion model). For irregularly shaped distributions we explored two different directions of contraction because distinct clines could influence results. The indexes were then validated exploring the behavior of values calculated at seven stages along the contraction process in these simulated scenarios (Supplementary material Fig. A3).

### *Regression analyses*

We defined regression models to estimate the probability of extirpation of an area based on two of the previously defined variables (*Border* and *Land use*) and the percentage of contraction (*Contraction*). For this approach we excluded distributions (ranges and fragments) with <10% or >90% contraction (Supplementary material, Appendix 1, Tables A1 and A2) because at early and late stages of contraction stochastic noise may confound existing patterns (Yackulic et al. 2011). Under the demographic model, the probability of extirpation should continuously decrease with the distance to the border independently of the percentage of contraction. Thus, the probability of extirpation of an area could be simply defined by the variable *Border* (*Mod\_Demographic*, Table 1). A key prediction of the contagion null model is that there is directionality in contraction,

but the angle of contraction is a relative concept that compares extirpated and current areas and thus, cannot be estimated for completely extirpated or current areas. Instead, we evaluated another prediction of this null model, namely that the effect of distance to the border on the probability of extirpation depends on the percentage of contraction. We modeled this prediction using an interaction term between the variables *Border* and *Contraction* (*Mod\_Contagion*, Table 1). Finally, under the refuge null model, the probability of extirpation should simply depend on the human land use intensity, which is represented by the variable *Land use* (*Mod\_Refuge*, Table 1). For each of the analysis scales (range and fragment) we fitted generalized linear mixed regression models (GLMM) with family binomial and a logit link using the function *glmer* from the *lme4* package in R (R Development Core Team 2013). All models included taxonomic class, order, family, and genus as random factors to control for evolutionary non-independence of the observations. We compared models using an information theoretic approach based on Akaike Information Criterion, AIC (Burnham and Anderson 2002).

Finally, we explored the possibility that the multiple processes postulated by these null models may occur simultaneously. We fitted two additional models that combine predictions from compatible null models. *Combined\_1* modelled the probability of extirpation considering both *Land use* and *Border*, *Combined\_2* included *Land use* and allowed for the interaction of *Border* with *Contraction* (Table 1).

## RESULTS

We analyzed spatial data for 386 species (374 species at range scale and 213 at fragment scale) which represent ~1.6% of the terrestrial vertebrates listed by the IUCN. The studied distribution ranges and fragments have widely variable areas, with an observed mean percentage of contraction of 41% for complete ranges and 51% for

fragments (Supplementary material Appendix 2, Tables A3 and A4 and Fig. A1). Distribution ranges are often fragmented with a mean of 6.7 fragments per historical range.

Validation of the indexes showed that as expected, when contraction was simulated following the demographic model, *Centrality Index* values decreased and *Directionality Index* values were generally close to 0 (although for irregular shapes values showed a small increase at high contraction stages). When contraction was simulated following a cline (as proposed by the contagion model), we detected the predicted shift in the *Centrality Index* and values for the *Directionality Index* generally close to 1.

Empirical estimates of the three indices did not identify a single best-supported null model at the range or fragment scale (Fig. 3). *Centrality Index* values show a tendency to change with the percentage of contraction as predicted by the contagion null model. However, *Directionality Index* values show no support for either the contagion or demographic models. The *Land use Index* suggests extirpation has been more likely in humanized areas as predicted by the refuge null model (median values are consistently above 1; Fig. 3). However, in many cases current areas are more humanized than those extirpated. Results were broadly consistent among taxonomic classes (Supplementary material, Appendix 2, Fig. A4).

Results from the regression analyses at both scales also failed to clearly identify a single best null model. At the range scale, both the refuge (*Mod\_Refuge*) and the contagion (*Mod\_Contagion*) null models received support; whereas at the fragment scale the only supported model was *Mod\_Refuge* (Table 1). Although overall the refuge null model received greater support compared to other null models, results at both range and fragment scales revealed that either of the combined models represents a great

improvement (based on AIC) over models based on the unifactorial null models (Table 1). At least for the available data, multiple processes appear to best explain the general patterns of contraction among terrestrial vertebrates.

At the range scale *Combined\_2* was the only supported model (Table 1), which describes the probability of extirpation as positively correlated with human use (*Land use*) and identifies a contraction-dependent effect of distance to the border. In particular, at early stages of contraction (up to ~60% contraction, obtained when the  $\partial\text{Probability of Extirpation}/\partial\text{Border}$  is equal to zero) areas near the border are more likely to be extirpated whereas at later stages the pattern is reversed (Fig. 4a). At the fragment scale, both combined models were supported (being within 2 AIC units of each other, Table 1) and show a positive association between the probability of extirpation and *Land use*, with the best supported model, *Combined\_2*, additionally supports an interaction between *Border* with *Contraction* with extirpation being generally more likely near the border, but with a weakening effect as contraction advances. In this model, extirpation only becomes more likely near the center outside the range of data values used to fit the model (approximately >98% contraction, obtained when the  $\partial\text{Probability of Extirpation}/\partial\text{Border}$  is equal to zero. Data used to fit the models exclude fragments with <10% or >90% contraction). The simpler supported model (*Combined\_1*) does not include an interaction term and suggests that extirpation is consistently more likely near the border (Figs. 4b and 4c). Thus, at the fragment scale, and considering both supported models we interpret the results as that in the early stages of contraction areas close to the border have higher probability of extirpation than central areas. However, this difference between border and central areas may weaken as contraction progresses. Separate analyses for data rasterized at different resolutions offered results consistent with these analyses (Supplementary material, Appendix 2, Table A9)

289

## 290 **DISCUSSION**

291 The three main null models of range contraction proposed to date make diverse  
292 predictions derived from their theoretical underpinnings. Our evaluation using global  
293 spatial data for terrestrial vertebrates reveals that none of these null models is  
294 sufficiently general to describe contraction range patterns. Even though in the majority  
295 of species extirpated areas are more likely to be heavily humanized, as predicted by the  
296 refuge null model, we also find support for models that incorporate two distinct  
297 mechanisms that likely act together. In addition, the relative position within a range also  
298 appears to influence extirpation probability (independently of human use). For many of  
299 the studied species, extirpation is more likely near the border during early stages of  
300 contraction but during the final stages of contraction extirpation becomes more likely in  
301 central areas, as proposed by the contagion null model. Yet, we also find support for the  
302 demographic model which postulates that the probability of extirpation is always higher  
303 near the border. Future research focused on the final stages of contraction would be  
304 necessary to disentangle these patterns. Nevertheless, our results show that contraction  
305 is better described by multi-process models that consider both human impacts and  
306 relative position, than by the three originally-proposed null models.

307

### 308 **Contraction and human land use**

309 We find that human use is probably the best single predictor of extirpation probability,  
310 as previously suggested by Yackulic et al. (2011). The key role of human land use  
311 changes in species extinction has been proposed by previous studies that identified  
312 habitat loss due to human land use as the main threat for diverse vertebrate groups  
313 (González-Suárez and Revilla 2014, Hayward 2011, Pekin and Pijanowski 2012,

Schipper et al. 2008). In our study, we find that indeed greater extirpation risk is generally associated with more humanized areas. However, a correlation between human use and extirpation does not imply a direct causal relationship. Other factors, such as the presence of invasive species or climate change, could be spatially correlated with human uses leading to similar patterns of contraction (Franco et al. 2006, Thomas et al. 2006). The potential role of these other factors could be explored considering our new proposed baseline that accounts for relative position and human impacts.

Although extirpations are generally more common in humanized areas, some species persist within these regions. Distinct patterns may be due to intrinsic responses; some species are less sensitive to human impacts than others (Maklakov et al. 2011), and some even benefit from humanized conditions (Maclean et al. 2011). Additionally, extirpation may be determined by other drivers of extinction with different spatial configurations (Clavero et al. 2009, González-Suárez et al. 2013, González-Suárez and Revilla 2014, Thomas et al. 2006). A caveat of our approach is that our data reflect only current human land uses, which may not correspond to the past uses potentially responsible for observed extirpations (Carvalho et al. 2013, Plieninger et al. 2006). It is not clear to us, however, how this could bias our results since we analyzed a large number of species at a global scale, and the progress of land use changes has been heterogeneous across the world. While land uses often intensify with time, the rates of intensification vary by area, and may affect species differently (Bregman et al. 2014, Gilroy et al. 2014). For example, in some areas of Europe and North America there has been a reversal toward more natural uses as agricultural land has been abandoned, but this reversal has not occurred in other areas (Gellrich et al. 2007, MacDonald et al. 2000, Mottet et al. 2006, Strijker 2005). Future studies would be necessary to address the temporal aspect of land use changes; however, human activities and land use are still



likely to be key factors driving range contraction. In fact, they may well play an even more important and complex role than identified here, e.g., areas with intense agricultural uses have a greater impact than agri-environmental management areas (Carvalho et al. 2013, Franco et al. 2006).

### **Contraction and relative position within the range: different patterns at different scales**

In addition to the importance of human land use, our analyses show that the relative position of an area also influences its probability of extirpation (Brown 1995, Channell and Lomolino 2000a, Channell and Lomolino 2000b, Lawton 1993). At the range scale our results indicate that the probability of extirpation near the border (or the center) depends on the contraction stage. This pattern can be caused by directional threats as proposed by Channell & Lomolino (2000a, 2000b). For example, climate change can create latitudinal and altitudinal clines (Parmesan 1996, Parmesan and Yohe 2003). However, there are alternative mechanisms that can also lead to this observed pattern. Climatic and biotic factors generally define range limits (Araújo and Rozenfeld 2014), but some boundaries are due to abrupt ecosystem changes or physical barriers, such as mountain chains or the transition from land to ocean. In these cases, border areas may actually represent optimal habitat and thus, be the most populated (Caughley et al. 1988, Gaston 2003, Sagarin and Gaines 2002). When optimal habitat occurs in a range border, a directional pattern of contraction could simply occur due to intrinsic population dynamics, as less dense populations are more likely to go extinct.

At the fragment scale we found support for two apparently contrasting models. The simplest model predicts that the probability of extirpation is always higher near the border, while the best model suggests that the probability of extirpation near the border

depends on the contraction level. However, the predicted shift from higher extirpation risk near the border to higher near the center occurs at the very final stages of contraction (which lay beyond the range of values analyzed, >90% contraction). In comparison, at the range scale this shift is predicted at ~60% contraction. Therefore, we interpret these results as supporting a higher probability of extirpation near fragment borders in early stages with a potential weakening of this effect as contraction progresses.

There are various possible reasons that could explain the discrepancy in the results between range and fragment scales. First, different factors and process influence dynamics at different scales, e.g., climate acts at broader scale while biotic interactions are more relevant locally (Araújo and Rozenfeld 2014, Pearson and Dawson 2003, Whittaker et al. 2001). Second, the meaning and identification of relative positions in complexly shaped distributions is complicated and this may confound results. For example, the border area in a fragment located near other fragments has a greater probability of receiving migrants than a “true border”, and thus, could have a lower probability of extirpation. Null models are commonly defined based on idealized distributions that largely fail to represent reality. Most species distributions are complex, often formed by multiple fragments with different shapes that change over time (Gaston 2003, Wilson et al. 2004). To study range dynamics we need to embrace this complexity, considering all types of ranges and not only those that conform to some theoretical or idealized depictions. Importantly, as shown here, we must evaluate predictions at different scales because results and inferences may differ (Thomas *et. al.* 2008).

**A new baseline to understand range contraction: multifactorial null models**

Earlier null models of range contraction have focused on single processes –basic population rules and simple threat dynamics (Brown and Kodricbrown 1977, Brown 1995, Channell and Lomolino 2000a, Channell and Lomolino 2000b, Lawton 1993). Here we show that these null models are not adequate baselines, at least for terrestrial vertebrates. Species persistence may be influenced by multiple external threats and intrinsic processes (González-Suárez et al. 2013, Yackulic et al. 2011). To partly account for this complexity, Yackulic *et al.* (2011) proposed multifactorial models (including biome, human impacts, and relative position) to explain range contraction in large mammals. Here, we generalized the importance of multifactorial models for a wide range of terrestrial vertebrates.

Understanding range contraction is important for conservation and management, particularly if we hope to accurately predict future range changes and assess the effects of new threats (Newbold et al. 2014, Peters et al. 2014, Selwood et al. 2014, Stanton et al. 2014, Thomas et al. 2004, Thomas et al. 2011). Our global study based on data from four different groups of vertebrates reveals the need to develop more realistic null models to use as baselines. Without departing from the objective of simplicity, we propose to combine simple key elements already identified as relevant to define new multi-process null models of range contraction. We realize that data at this scale could have their own limitations, but we feel that these models can offer a more realistic baseline to evaluate the role of additional factors, such as the effect of different types of range borders, the role of environmental conditions, additional human and natural threats, as well as how intrinsic species' traits influence contraction range dynamics.

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## REFERENCES

- Araújo, M. B. and Rozenfeld, A. 2014. The geographic scaling of biotic interactions. - *Ecography* 37: 406–415.
- Bregman, T. P. et al. 2014. Global patterns and predictors of bird species responses to forest fragmentation: implications for ecosystem function and conservation. - *Biol. Conserv.*
- Brown, J. H. 1971. Mammals on mountaintops: nonequilibrium insular biogeography. - *Am. Nat.* 467-478.
- Brown, J. H. and Kodricbrown, A. 1977. Turnover rates in insular biogeography: effect of immigration on extinction<sup>1</sup>. - *Ecology* 58: 445-449.
- Brown, J. H. 1995. *Macroecology*. - University of Chicago Press, Chicago.

438 Burnham, K. P. and Anderson, D. R. 2002. Model Selection and Multimodel Inference:  
 439 A Practical Information-Theoretic Approach. - Springer, New York.  
 440 Cahill, A. E. et al. 2012. How does climate change cause extinction? - Proc. R. Soc.  
 441 Lond., Ser. B: Biol. Sci. 280: 1-9.  
 442 Carvalheiro, L. G. et al. 2013. Species richness declines and biotic homogenisation have  
 443 slowed down for NW-European pollinators and plants. - Ecol. Lett. 16: 870-878.  
 444 Caughley, G. et al. 1988. The edge of the range. - The Journal of Animal Ecology  
 445 Ceballos, G. and Ehrlich, P. R. 2002. Mammal population losses and the extinction  
 446 crisis. - Science 296: 904-907.  
 447 Channell, R. and Lomolino, M. V. 2000a. Dynamic biogeography and conservation of  
 448 endangered species. - Nature 403: 84-86.  
 449 Channell, R. and Lomolino, M. V. 2000b. Trajectories to extinction: spatial dynamics of  
 450 the contraction of geographical ranges. - J. Biogeogr. 27: 169-179.  
 451 Clavero, M. et al. 2009. Prominent role of invasive species in avian biodiversity loss. -  
 452 Biol. Conserv. 142: 2043–2049.  
 453 David, H. R. et al. 2003. Estimates of minimum viable population sizes for vertebrates  
 454 and factors influencing those estimates. - Biol. Conserv. 113: 2334.  
 455 Davis, A. J. et al. 1998. Making mistakes when predicting shifts in species range in  
 456 response to global warming. - Nature 391: 783-786.  
 457 Fisher, D. O. 2011. Trajectories from extinction: where are missing mammals  
 458 rediscovered? - Global Ecol. Biogeogr. 20: 415-425.  
 459 Franco, A. M. A. et al. 2006. Impacts of climate warming and habitat loss on extinctions  
 460 at species' low-latitude range boundaries. - Global Change Biol. 12: 1545-1553.  
 461 Gaston, K. J. 2003. The Structure and Dynamics of Geographic Ranges. - Oxford  
 462 University Press, Oxford.

463 Gellrich, M. et al. 2007. Agricultural land abandonment and natural forest re-growth in  
 464 the Swiss mountains: A spatially explicit economic analysis. - *Agric., Ecosyst. Environ.*  
 465 118: 93-108.  
 466 Gilroy, J. J. et al. 2014. Effect of scale on trait predictors of species responses to  
 467 agriculture. - *Conserv. Biol.*  
 468 González-Suárez, M. et al. 2013. Which intrinsic traits predict vulnerability to  
 469 extinction depends on the actual threatening processes. - *Ecosphere* 4 (6): 76.  
 470 González-Suárez, M. and Revilla, E. 2014. Generalized drivers in the Mammalian  
 471 endangerment process. - *PloS ONE* 9: e90292.  
 472 Gotelli, N. J. 2001. Research frontiers in null model analysis. - *Global Ecol. Biogeogr.*  
 473 10: 337-343.  
 474 Hanski, I. 1998. Metapopulation dynamics. - *Nature* 396: 41-49.  
 475 Hanski, I. and Ovaskainen, O. 2000. The metapopulation capacity of a fragmented  
 476 landscape. - *Nature* 404: 755-758.  
 477 Hayward, M. W. 2011. Using the IUCN Red List to determine effective conservation  
 478 strategies. - *Biodivers. Conserv.* 20: 2563–2573.  
 479 Hemerik, L. et al. 2006. The eclipse of species ranges. - *Acta Biotheor.* 54: 255-266.  
 480 Hoffmann, M. et al. 2010. The Impact of Conservation on the Status of the World's  
 481 Vertebrates. - *Science* 330: 1503-1509.  
 482 International Union for Conservation of Nature 2010. IUCN red list of threaten species  
 483 Version 2010.4. <http://www.iucnredlist.org/> (accessed February 08, 2010).  
 484 Jones, H. L. and Diamond, J. M. 1976. Short-time-base studies of turnover in breeding  
 485 bird populations on the California Channel Islands. - *Condor*  
 486 Laliberte, A. S. and Ripple, W. J. 2004. Range contractions of North American  
 487 carnivores and ungulates. - *Bioscience* 54: 123-138.

488 Lawton, J. H. 1993. Range, population abundance and conservation. - Trends Ecol.  
 489 Evol.  
 490 Li, X. et al. 2015. Human impact and climate cooling caused range contraction of large  
 491 mammals in China over the past two millennia. - Ecography 38: 74–82.  
 492 MacArthur, R. H. and Wilson, E. O. 1967. The Theory of Island Biogeography. -  
 493 Princeton University Press, Princeton, N.J.  
 494 MacDonald, D. et al. 2000. Agricultural abandonment in mountain areas of Europe:  
 495 Environmental consequences and policy response. - J. Environ. Manage. 59: 47-69.  
 496 Maclean, I. et al. 2011. Predicting changes in the abundance of African wetland birds by  
 497 incorporating abundance– occupancy relationships into habitat association models. -  
 498 Divers. Distrib. 17: 480-490.  
 499 Maklakov, A. A. et al. 2011. Brains and the city: big-brained passerine birds succeed in  
 500 urban environments. - Biol. Lett. 7: 730–732.  
 501 Mottet, A. et al. 2006. Agricultural land-use change and its drivers in mountain  
 502 landscapes: A case study in the Pyrenees. - Agric., Ecosyst. Environ. 114: 296-310.  
 503 Newbold, T. et al. 2014. Functional traits, land-use change and the structure of present  
 504 and future bird communities in tropical forests. - Global Ecol. Biogeogr. 23: 1073-1084.  
 505 Oak Ridge National Laboratory Distributed Active Archive Center 2010. MODIS  
 506 (MCD12Q1) Land Cover Product <http://webmap.ornl.gov> ORNL DAAC, Oak Ridge,  
 507 Tennessee, USA. Accessed September, 2013.  
 508 Parmesan, C. 1996. Climate and species' range. - Nature 382: 765-766.  
 509 Parmesan, C. and Yohe, G. 2003. A globally coherent fingerprint of climate change  
 510 impacts across natural systems. - Nature 421: 37-42.

511 Pearson, R. G. and Dawson, T. P. 2003. Predicting the impacts of climate change on the  
 512 distribution of species: are bioclimate envelope models useful? - *Global Ecol. Biogeogr.*  
 513 12: 361-371.

514 Pekin, B. K. and Pijanowski, B. C. 2012. Global land use intensity and the  
 515 endangerment status of mammal species. - *Divers. Distrib.* 18: 909-918.

516 Peters, H. et al. 2014. Identifying species at extinction risk using global models of  
 517 anthropogenic impact. - *Global Change Biol.* doi: 10.1111/gcb.12749.

518 Pimm, S. L. et al. 1988. ON THE RISK OF EXTINCTION. - *Am. Nat.* 132: 757-785.

519 Plieninger, T. et al. 2006. Traditional land-use and nature conservation in European  
 520 rural landscapes. - *Environ. Sci. Policy* 9: 317-321.

521 Pomara, L. Y. et al. 2014. Demographic consequences of climate change and land cover  
 522 help explain a history of extirpations and range contraction in a declining snake species.  
 523 - *Global Change Biol.* 20: 2087–2099.

524 R Development Core Team 2013. R: A language and environment for statistical  
 525 computing. R Foundation for Statistical Computing, Vienna, Austria.

526 Safi, K. and Pettorelli, N. 2010. Phylogenetic, spatial and environmental components of  
 527 extinction risk in carnivores. - *Global Ecol. Biogeogr.* 19: 352-362.

528 Sagarin, R. D. and Gaines, S. D. 2002. The 'abundant centre' distribution: to what extent  
 529 is it a biogeographical rule? - *Ecol. Lett.* 5: 137-147.

530 Sagarin, R. D. et al. 2006. Moving beyond assumptions to understand abundance  
 531 distributions across the ranges of species. - *Trends Ecol. Evol.* 21: 524-530.

532 Schipper, J. et al. 2008. The status of the world's land and marine mammals: diversity,  
 533 threat, and knowledge. - *Science* 322: 225-230.



534 Selwood, K. E. et al. 2014. The effects of climate change and land-use change on  
 535 demographic rates and population viability. - *Biol. Rev. Camb. Philos. Soc.* doi:  
 536 10.1111/brv.12136.

537 Stanton, J. C. et al. 2014. Warning times for species extinctions due to climate change. -  
 538 *Global Change Biol.* doi: 10.1111/gcb.12721.

539 Strijker, D. 2005. Marginal lands in Europe—causes of decline. - *Basic Appl. Ecol.* 6:  
 540 99-106.

541 Thomas, C. et al. 2006. Range retractions and extinction in the face of climate warming.  
 542 - *Trends Ecol. Evol.* 21: 415-416.

543 Thomas, C. D. et al. 2004. Extinction risk from climate change. - *Nature* 427: 145-148.

544 Thomas, C. D. et al. 2008. Where within a geographical range do species survive best?  
 545 A matter of scale. - *Insect Conserv. Divers.* 1: 2-8.

546 Thomas, C. D. et al. 2011. A framework for assessing threats and benefits to species  
 547 responding to climate change. - *Methods Ecol. Evol.* 2: 125-142.

548 Turvey, S. T. et al. 2015. Historical data as a baseline for conservation: reconstructing  
 549 long-term faunal extinction dynamics in Late Imperial–modern China. - *Proc. R. Soc. B*

550 Whittaker, R., J. et al. 2001. Scale and species richness: towards a general, hierarchical  
 551 theory of species diversity. - *J. Biogeogr.* 28: 453-470.

552 Wilson, R. J. et al. 2004. Spatial patterns in species distributions reveal biodiversity  
 553 change. - *Nature* 432: 393-396.

554 Yackulic, C. B. et al. 2011. Anthropogenic and environmental drivers of modern range  
 555 loss in large mammals. - *Proc. Natl. Acad. Sci. USA* 108: 4024-4029.

556 Zar, J. H. 1999. *Biostatistical analysis*, 4/e. - Pearson Education India.

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558 **SUPPLEMENTARY MATERIAL**

- 559    Supplementary material (Appendix EXXXXXX at <[www.oikosoffice.lu.se/appendix](http://www.oikosoffice.lu.se/appendix)>).
- 560    Appendices 1-2.

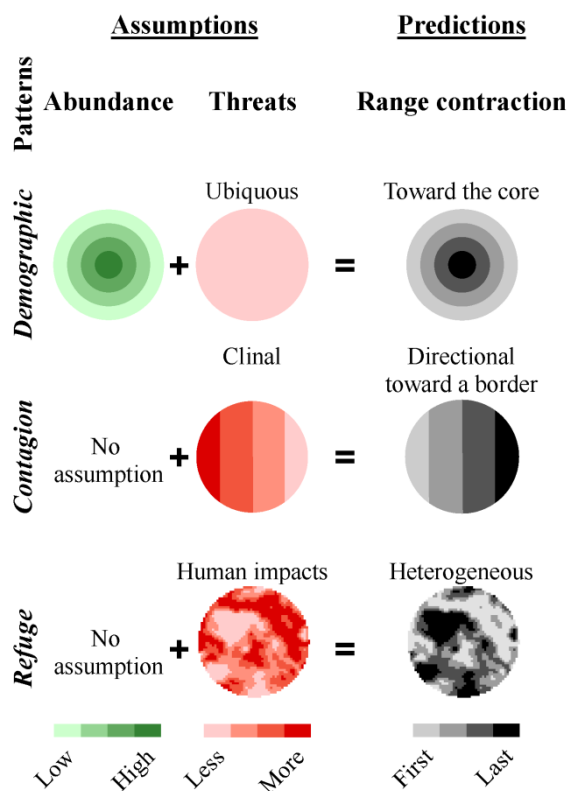
561    **TABLES**

562    **Table 1.** Results from the regression analyses based on regression models (GLMM) to evaluate the three main null models of range contraction  
563    (demographic, contagion and refuge) and two combined models that incorporate multiple processes. *Combined\_1* proposes that the probability of  
564    extirpation of an area is determined by the proportion of human use in the area (variable *Land use*) and the distance to the historical border  
565    (variable *Border*). *Combined\_2* proposes that the probability of extirpation depends on *Land use* and the interaction of *Border* and *Contraction*  
566    (reflecting the expectation that as range contraction progresses the risk associated with being near the border changes). All models were fitted at  
567    two scales: complete historical range and historical fragment. We report model coefficients (best estimates and their SE), AIC,  $\Delta$ AIC (difference  
568    in AIC with the best model comparing all five models), and  $\Delta$ AIC<sub>sm</sub> (difference in AIC comparing only the three models derived from the main  
569    proposed null models). Dashes indicate variables not included in the model.

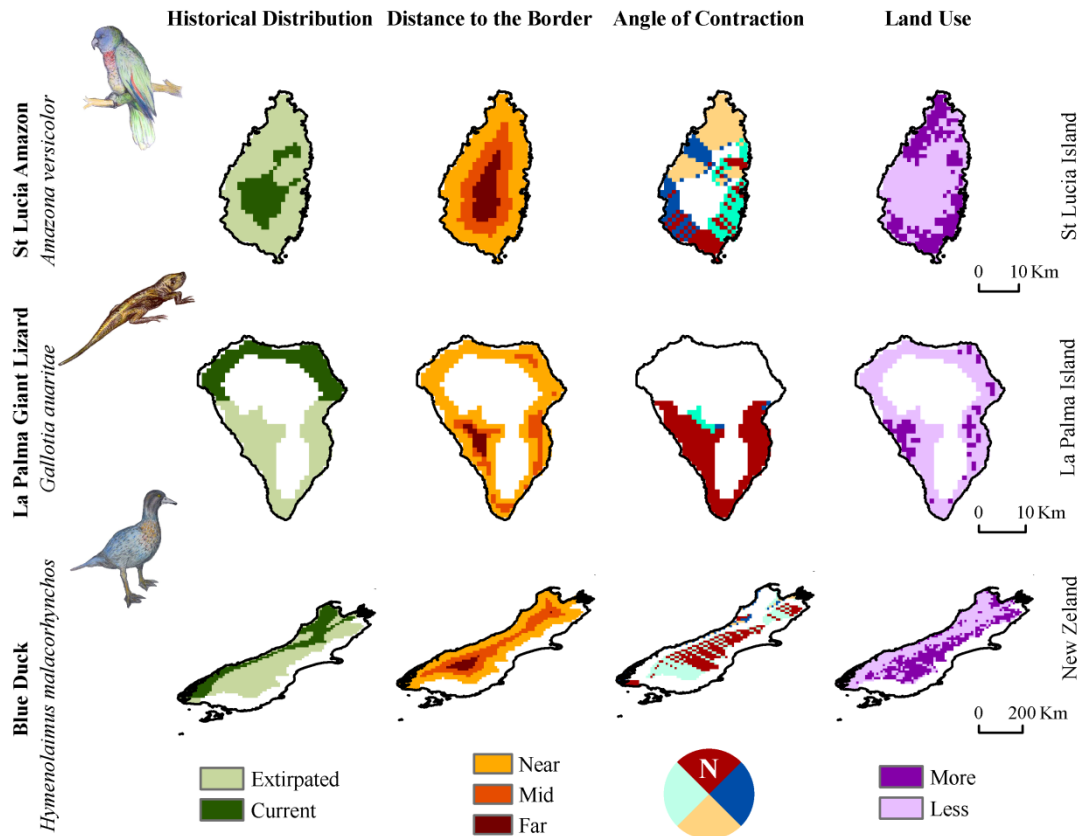
Model	Coefficients				Model comparison		
	<i>Land use</i>	<i>Border</i>	<i>Contraction</i>	<i>Border*Contraction</i>	<i>AIC</i>	$\Delta AIC$	$\Delta AIC_{sm}$
<b>Range scale (N=457, 229 species)</b>							
<i>Combined_2</i>	2.13 (0.466)*	-9.74 (2.145)*	-2.66 (0.688)*	15.86 (3.699)*	605.21	0.00	
<i>Combined_1</i>	2.03 (0.443)*	-1.78 (0.919) †	-	-	621.33	16.13	
<i>Mod_Refuge</i>	2.02 (0.441)*	-	-	-	623.15	17.94	0.00
<i>Mod_Contagion</i>	-	-9.81 (2.110)*	-2.23 (0.664)*	15.74 (3.650)*	625.49	20.28	2.34
<i>Mod_Demographic</i>	-	-1.74 (0.887) †	-	-	641.64	36.43	18.49
<b>Fragment scale (N=362, 142 species)</b>							
<i>Combined_2</i>	2.73 (0.541)*	-9.15 (2.497)*	-2.03 (0.977)*	9.35 (4.131)*	468.09	0.00	
<i>Combined_1</i>	2.62 (0.527)*	-4.16 (1.008)*	-	-	469.35	1.26	
<i>Mod_Refuge</i>	2.57 (0.514)*	-	-	-	486.24	18.14	0.00
<i>Mod_Contagion</i>	-	-8.30 (2.430)*	-1.22 (0.927)	7.65 (3.975) †	494.30	26.21	8.06
<i>Mod_Demographic</i>	-	-3.98 (0.952)*	-	-	494.72	26.62	8.48

570 \*P < 0.05; †P < 0.10

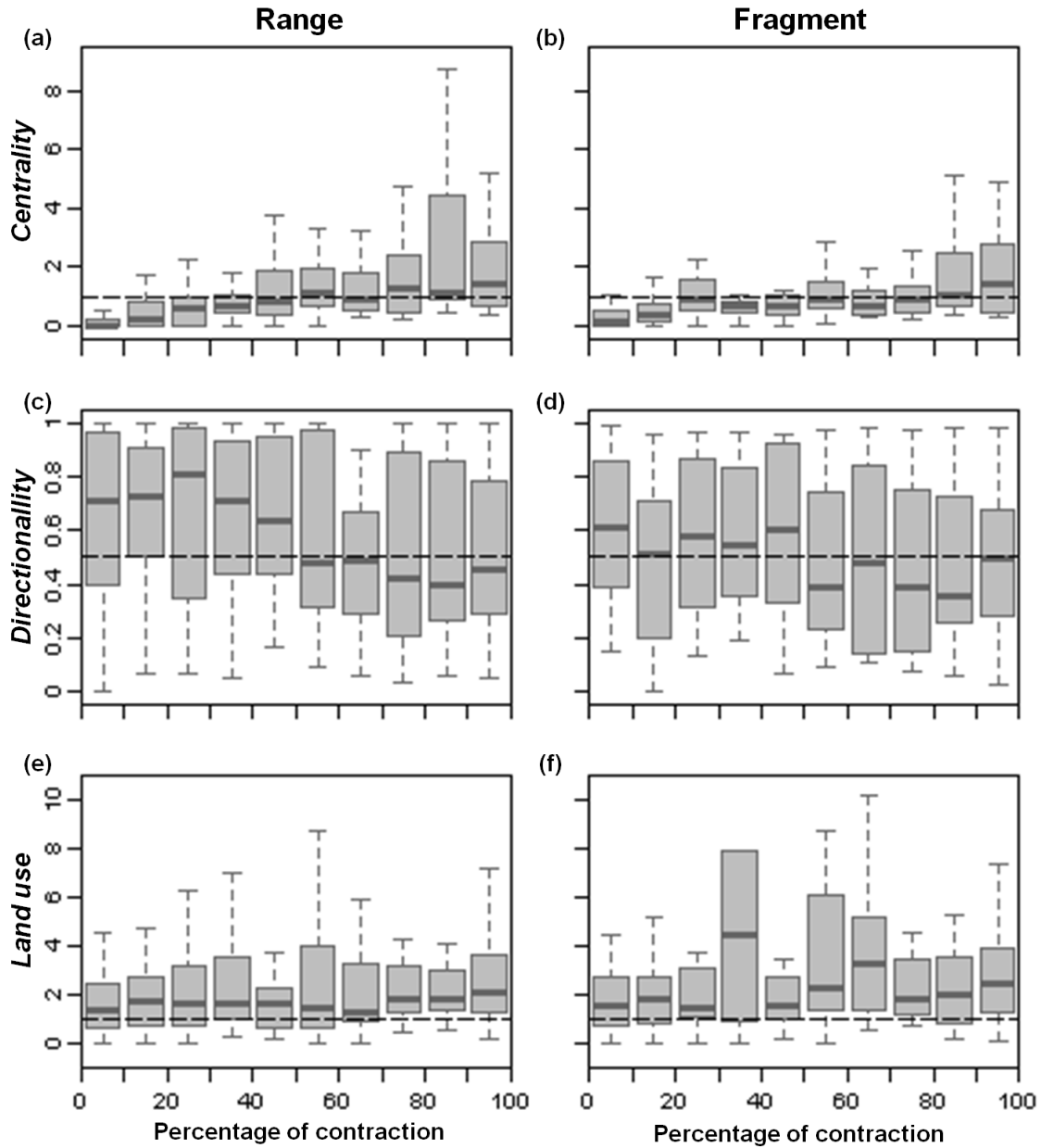
**FIGURES**



**Figure 1.** Assumptions and predicted range contraction patterns for each of the three null models. The demographic null model assumes higher density in the center of the range and a ubiquitous threat pattern. As a result, contractions are predicted to occur toward the core in multiple directions. The contagion null model assumes that threats are distributed in a cline resulting in a directional contraction along this cline. The refuge null model assumes that the extirpation is determined by human land use and predicts a heterogeneous range contraction pattern with less used areas being less likely to become extirpated.



**Figure 2.** Examples of the three variables defined to represent the key predictors of the three null models: Distance to the Border (*Border*): average distance to border from each cell; *Angle of contraction*: geodetic angle of contraction (from each extirpated cell to the closest current cell), and Human Use (*Land use*): proportion of human use in the cell. Examples represent the Saint Lucia amazon (*Amazona versicolor*) which illustrates the pattern of contraction predicted by the demographic null model (also partly congruent with the refuge null model); the La Palma giant lizard (*Gallotia auaritae*) illustrates contraction from a border to the opposite border in a unique direction as predicted by the contagion null model (and is also partly congruent with the refuge null model); and the blue duck (*Hymenolaimus malacorhynchos*) which adjusts to the refuge null model prediction.

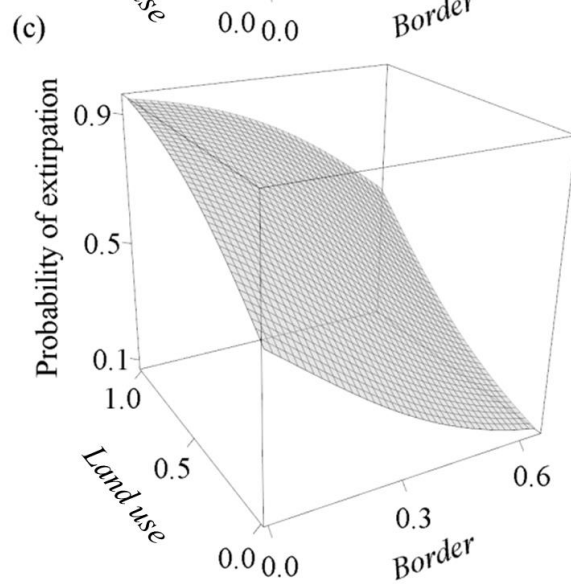
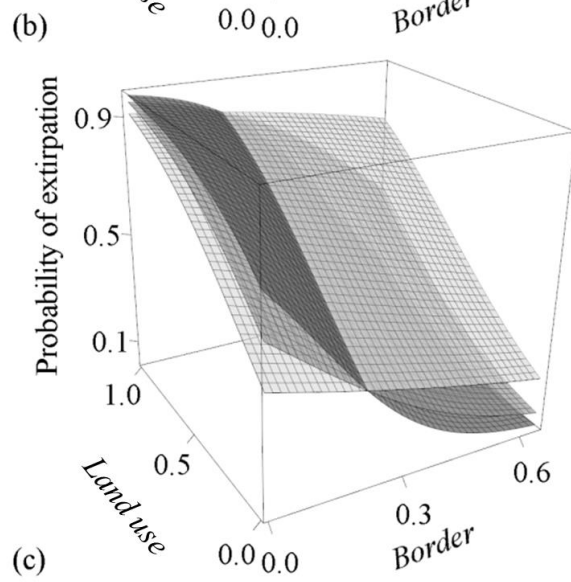
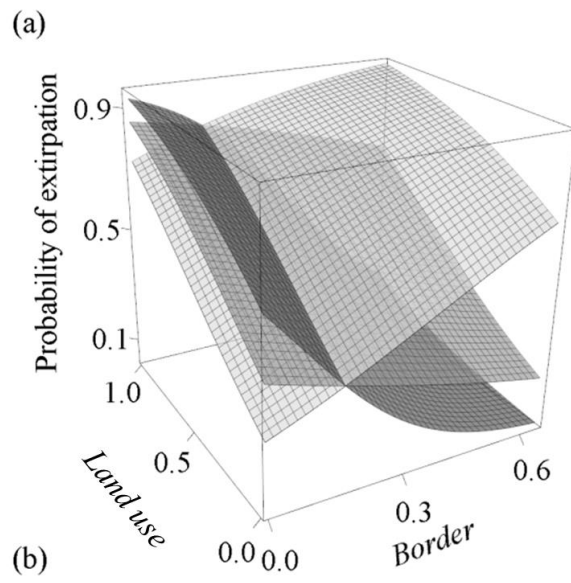


**Figure 3.** The distribution of three indices at the range (a, c, e) and fragment scale (b, d, f). For initial stages of contraction (< 50% contraction) both demographic and contagion null model predict *Centrality Index* < 1. For higher stages of contraction (> 50% contraction) *Centrality Index* < 1 supports the demographic null model while *Centrality Index* > 1 supports the contagion null model (a, b). *Directionality Index* close to 0 is predicted by the demographic null model, whereas values close to 1 support the contagion null model (c, d). *Land use Index* > 1 is predicted by the refuge null model (e, f). Ends of the whiskers represent the lowest datum still within the 1.5 interquartile

604 range (IQR) of the lower quartile, and the highest datum still within the 1.5 IQR of the  
605 upper quartile (Tukey boxplot).

606





**Figure 4.** Predictions of the supported regression models explaining probability of extirpation of an area as a function of its distance to the historical border (*Border*) and its human land use (*Land use*) with a possible interaction of *Land use* and the percentage of contraction (*Contraction*). At the range scale, panel (a), Model *Combined\_2* (including the interaction) was the single supported model (Table 1). At the fragment scale both Model *Combined\_2* (b) and Model *Combined\_1* (c, no interaction) were supported. To visualize the effect of the interaction between *Border* and *Contraction* (a, b), we represent predictions at three levels of contraction: 20% in darker grey, 50% in medium dark grey, and 80% in light grey.